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RAW SEQUENCE LISTING TRADEMAN ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/8/2,315

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510: FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or-other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1903-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/8/2/5/5
ATTN: NEW RULES CASES	: Please disregard english "Alpha" headers, which were inserted by PTO softwari
Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6Patentin 2.0 "bug"	A "bug" in Patentin version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, Patentin would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7 Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number
•	000
9Use of n's or X2a's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10 V Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220> <223> section is required when <213> response is Unknown of is Artificial Sequence.
11Use of <220>	Sequence(s)
Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>arnino acid</u>

AMC - Biotechnology Systems Branch - 09/09/2003





IFWO

RAW SEQUENCE LISTING

DATE: 04/06/2004

PATENT APPLICATION: US/10/812,315

TIME: 10:51:23

Input Set : A:\Sequence Listing 81000.txt Output Set: N:\CRF4\04062004\J812315.raw

```
4 <110> APPLICANT: Degussa AG
      6 <120> TITLE OF INVENTION: A process for producing L-amino acids using strains of the
              Enterobacteriaceae family
      9 <130> FILE REFERENCE: 020481 BT
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/812,315
C--> 11 <141> CURRENT FILING DATE: 2004-03-30
     11 <160> NUMBER OF SEQ ID NOS: 4
    13 <170> SOFTWARE: PatentIn version 3.1
    15 <210> SEQ ID NO: 1
    16 <211> LENGTH: 32
    17 <212> TYPE: DNA
                                                 Does Not Comply
    18 <213> ORGANISM: (Synthetic sequence
    21 <220> FEATURE:
                                                        Corrected Diskette Needed
W--> 22 <221> NAME/KEY: Primer
                                                        in Ulynknown
    23 <222> LOCATION: (1)..(32)
    24 <223> OTHER INFORMATION: galP1
    27 <400> SEQUENCE: 1
    28 cacaatctag ataaaccata ttggagggca tc
    30 <210> SEQ ID NO: 2
    31 <211> LENGTH: 25
     32 <212> TYPE: DNA
     33 <213> ORGANISM: (Synthetic sequence
     36 <220> FEATURE:
  -> 37 <221> NAME/KEY: Primer
    38 <222> LOCATION: (1)..(25)
     39 <223> OTHER INFORMATION: galP2
    42 <400> SEQUENCE: 2
     43 gggaggaage ttggggagat taate
     45 <210> SEQ ID NO: 3
    46 <211> LENGTH: 1446
     47 <212> TYPE: DNA
     48 <213> ORGANISM: Escherichia coli
    51 <220> FEATURE:
W--> 52 <221> NAME/KEY: DNA fragment
    53 <222> LOCATION: (1)..(1446)
    54 <223> OTHER INFORMATION: PCR product
    57 <220> FEATURE:
     58 <221> NAME/KEY: CDS
     59 <222> LOCATION: (33)..(1427)
     60 <223> OTHER INFORMATION: galP coding region
W--> 61 <400> SEQUENCE: 3
     62 cacaatctag ataaaccata ttggagggca to atg cot gac got aaa aaa cag
```

Met Pro Asp Ala Lys Lys Gln

63

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/812,315 TIME: 10:51:23

DATE: 04/06/2004

Input Set : A:\Sequence Listing 81000.txt
Output Set: N:\CRF4\04062004\J812315.raw

														_				
64										1				5			101	
· 66	ggg	cgg	tca	aac	aag	gca	atg	acg	ttt	ttc	gtc	tgc	ttc	ctt	gcc	gct	101	
67	Gly	Arg	Ser	Asn	Lys	Ala	Met	Thr	Phe	Phe	Val	Cys		Leu	Ala	Ala		
68			10					15					20					
70	ctg	gcg	gga	tta	ctc	ttt	ggc	ctg	gat	atc	ggt	gta	att	gct	ggc	gca	149	
71	Leu	Ala	Gly	Leu	Leu	Phe	Gly	Leu	Asp	Ile	Gly	Val	Ile	Ala	Gly	Ala		
72		25					30					35						
74	ctg	ccg	ttt	att	gca	gat	gaa	ttc	cag	att	act	tcg	cac	acg	caa	gaa	197	
75	Leu	Pro	Phe	Ile	Ala	Asp	Glu	Phe	Gln	Ile	Thr	Ser	His	Thr	Gln	Glu		
76	40					45					50					55 _.		
78	tgg	gtc	gta	agc	tcc	ätg	atg	ttc	ggt	gcg	gca	gtc	ggt,	gcg	gtg	ggc	245	
79	Trp	Val	Val	Ser	Ser	Met	Met	Phe	Gly	Ala	Ala	Val	Gly	Ala	Val	Gly		
80	•				60				_	65					70			
	agc	aac	taa	ctc	tcc	ttt	aaa	ctc	aga	cqc	aaa	aag	agc	ctg	atg	atc	293	
83	Ser	Glv	Tro	Leu	Ser	Phe	Lvs	Leu	Glv	Arq	Lys	Lys	Ser	Leu	Met	Ile		
84	001	01	1.4	75			-,-		80		•	-		85				
	aac	aca	att	ttg	ttt	att	acc	aat	t.ca	cta	ttc	tct	aca	act	qcq	cca	341	
97	Clu	λla	Tla	Leu	Phe	Val	Ala	Glv	Ser	Leu	Phe	Ser	Ala	Ála	Ála	Pro		
88	Gry	PTG	90	Бец	1110			95	002				100					
		~++		gta	cta	a++	ctt		cac	att	cta	cta		cta	aca	ata	389	
90	aac	gict	Clu	y ca	Lou	Tla	Lon	Sor	Ara	Val	Len	Len	Glv	Len	Ala	Val		
	ASII	105	GIU	Val	ьęи	116	110	Ser	Arg	V (1.1	ыси	115	or,	200				
92				tct	+ ~ +			~~~	ota	+ 20	ctc		naa	att	aca	cca	437	
94	ggt	gtg	gcc	Ser	Tur	The T	y Ca	Dro	LLY	Tur	Tan	Sar	Glii	Tla	Ala	Pro		
		vai	ATG	ser	ıyı		Ald	PIÓ	Leu	1 7 1	130	261	GIU	110	niu	135		
	120					125		2+2	+ 0.0	2+4		C 3 C	tta	ata.	atc.		485	
98	gaa	aaa	att	cgt	ggc	agu	aty	TIA	Com	Mat	Tur	Cla	Lau	Mot	Tla	Thr	,00	
		гля	116	Arg			мес	116	Ser	145		GIII	neu	rie c	150			
100) ` .				140		مدند مد							- 200			533	
107	z ato	999	g ato	CLC	ggu	, gcc	. lai			yat	. acc	- 9C	Dha	. Car	e Tu	acc	333	
		e GT	A 116			Ala	a lyi	ret			3 1111	. MIC	1 E110	165		Thr		
104				155					160								581	
100	6 gg1	gca	a tgo	g cgc	: tgc	ato	CLO	ggt	gro	all	. dl(. all	. Dv	y yce	1 all	ttg	301	
		Ala			rrp) Met	ret			. 116	: 116	3 116	180		1 116	Leu		
108	3		170					175							_ `		629	
110) ct	gct	gatt	: ggt	gto	tto	tto	cti	g cca	gac	ago	CC	a egu	t t g	ם בנו	gcc	023	
11:	l Lei			e Gry	, val	Phe			Pro) AS	o ser			1 1 1 1) PHE	Ala		
11:		185					190					19					677	
114	a gco	c aaa	a cgo	cgt	ttt	gtt	. gat	. gcc	gaa	cgo	gto	cte	g cta	ı cgo	cti	g cgt	677	
11:	5 Ala	a Lys	s Arg	g Arg	l bye) Ala	a Glu	Arç			ı Lei	ı Arç	g rer	Arg		
	6 200					205					210	-				215	705	
118	3 gad	aco	c ago	c gcg	g gaa	gco	j aaa	cgc	c, gaa	ctç	g gat	: gaa	aato	cgt	: gaa	agt	725	
119	9 Ası	Th:	c Sei	. Ala	Glu	Ala	Lys	Arg	g Glu			o Glu	ı Ile	e Arg	g Gli	Ser	•	
120					220					225	-				230			
122	2 ttq	g cag	g gtt	: aaa	caç	gagt	ggc	: tgg	g gcg	, ct	; ttt	aaa	a gaç	g aac	c ago	aac	773	
123	3 Let	ı Glr	ı Val	Lys	Glr	ı Ser	: Gl	, Trp	o Ala	Let	ı Phe	E Lys	s Glu	ı Ası	ı Sei	Asn	•	
124				235					240					24				
120	6 tto	cgc	c cgc	geg	gto	tto	ctt	ggc	gta	cte	tto	caq	g gta	aatq	g cag	caa	821	
12	7 Phe	e Arc	Arc	, Ala	Va]	Phe	Let	Gly	/ Val	Leu	ı Let	ı Glı	n Val	l Met	: Glr	Gln		
-128		-,	250			•		255					260					

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/812,315

DATE: 04/06/2004 TIME: 10:51:23

Input Set : A:\Sequence Listing 81000.txt
Output Set: N:\CRF4\04062004\J812315.raw

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	131	Phe	Thr	Glv	Met	Asn	vаl	Ile	Met	Tyr	Tyr	Ala	Pro	Lys	Ile	Phe	Glu	
	132		265	1				270		-			275					
		cta		aat	tat	acc	aac	act	acc	qaq	caa	atq	tqq	ggg	acc	gtg	att	917
	135	Leu	Ala	Glv	Tyr	Thr	Asn	Thr	Thr	Glu	Gln	Met	Trp	Gly	Thr	Val	Ile	
		280		1	- , -		285					290	-	-			295	
			aac	cta	acc	aac		ctt	qcc	acc	ttt	atc	qca	atc	ggc	ctt	gtt	965
	139	Val	Glv	Len	Thr	Asn	Val	Leu	Ala	Thr	Phe	Ile	Āla	Ile	Ğĺy	Leu	Val	
	140		U 1	200		300					305			•		310		
		nac	cac	taa	gga		aaa	cca	aca	cta	acq	cta	aac	ttc	ctg	gtg	atg	1013
	143	Δsn	Ara	Trn	Gly	Ara	Lvs	Pro	Thr	Leu	Thr	Leu	Ğĺv	Phe	Leu	Val	Met	
	144	пор	****	112	315	****9	2,0			320			,		325			
		act	act	aac	atg	aac	ota	ctc	aat		atα	ata	cat	atc	aat	att	caċ	1061
	140	Ala	Δla	Glv	Met	Glv	Val	Len	Glv	Thr	Met	Met	His	Ile	Ğĺν	Ile	His	
	148	нта	AT O	330	1700		vui	БСС	335					340				
	150	+ <+	cca		gcg	can	tat	ttc		atc	acc	ato	cta		atg	ttt	att	1109
	151	Cor	Dro	Sor	Ala	Gln	Tur	Phe	Ala	Tle	Ala	Met	Leu	Leu	Met	Phe	Ile	•
	152	261	345	561	niu	0111	. , .	350					355					
		atc		+++	gcc	atro	ant		aat	cca	cta	att		ota	cta	tac	tcc	1157
	155	Val	Clu	Dha	Ala	Mot	Ser	Δla	Glv	Pro	Len	Tle	Trp	Val	Leu	Cvs	Ser	
		360	GLY	riie	n.a	1700	365	711.0	G1		200	370				- 2	375	
			2++	can	ccg	cta		aac	ćac	gat	+++	-	atc	acc	tac	tcc	act	1205
	150	Clu	Tla	Cln	Pro	Len	Lvs	610	Ara	Asn	Phe	Glv	Ile	Thr	Cvs	Ser	Thr	
	160	GIU	116	GTH	110	380	БУЗ	O.L.y	1119	тор	385	01,			-,-	390		
		~~~	200	220	tgg		acc	220	ato	atc		aac	aca	acq	ttc		acc	1253
	162	Ala	The	Acr	Trp	Tla	Ala	Acn	Mot	Tle	Val	Glu	Ala	Thr	Phe	Leu	Thr	
	164	нта	TIIT	VOII	395	116	VIG	non	1100	400	•	01,			405			
	166	2+4	ot c	220	acg	cta	aat	220	acc		acc	ttc	taa			aca	act	1301
	167	Mot	Lou	Aco	Thr	Len	Glu	Asn	Ala	Asn	Thr	Phe	Tro	Val	Tvr	Ala	Ala	•
	168	met	ren	410	1111	Deu	Gry	71.511	415	110.1				420	-,-			
		ota	220		ctg	+++	atc	cta		aca.	tta	taa	cta		cca	gaa	acc	1349
	171	Lou	Acn	yca Val	Leu	Dha	Tla	Leu	Len	Thr	Len	Trn	Len	Val	Pro	Glu	Thr	
	172	ьeu	425	vai	Deu	LIIC	116	430	БСС	****	ВСС		435					
		222		att	tcg	cta	as a		att	gaa	cat	aat		atσ	aaa	aat	cat	1397
•	175	Luc	ui.c	Val	Ser	Leu	Glu	Hie	Tle	Glu	Ara	Asn	Leu	Met	Lvs	Glv	Ara	
		440	nis	491	JCI	DCu	445			010		450			-,-		455	
			ota	cac	gaa	at a		act	cac	gat	taa		cocca	aaa e	cttc	ctcc		1446
					Glu						cuu						-	
	180	гу	Leu	Arg	GIU	460	СТУ	AIG	1113	1135					•			
		c216	) > CI	וז מי	O NO:													
					H: 40													
		_		PE:		<b>J</b> 1												
					ISM:	Feci	heri	-hia	col i									
					NCE:		CLIC	JIII	0011	_								
	100	Mot	D~C	ソクト	Ala	Lue	Lve	Gln	Glv	Ara	Ser	Asn	Lvs	Ala	Met	Thr	Phe	
			FIO	чэр	utq	r y	ny 3	7111	OLY.	*** 9	10		-10			15		
	190		1/-1	Cuc	Phe	J	מומ	ב ו ב	Leu	Δls		Len	Len	Phe	Gl v		Asp	
		rne	AGT	cys		ьец	uiq	nid	neu	25	OT A	±÷u	Lou		30			
	193	T1.	C1	1107	20 Ile	~ 1 ת	C1	7 l ~	ton		Phe	Tle	Δla	Asn		Phe	Gln	
	132	тте	сτλ	vai	116	HIG	GIÀ	wrd	reu	LIO	File	TIG	A14	nsp	O L U	2.10		

#### RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/812,315

DATE: 04/06/2004 TIME: 10:51:23

Input Set : A:\Sequence Listing 81000.txt
Output Set: N:\CRF4\04062004\J812315.raw

196			35			• •		40					45.			
198	Ile	Thr	Ser	His	Thr	Gln	Glu	Trp	Val	Val	Ser	Ser	Met	Met	Phe	Gly
199		50					55					60				
201	Ala	Ala	Val	Gly	Ala	Val	Gly	Ser	Gly	Trp	Leu	Ser	Phe	Lys	Leu	Gly
202	65					70					75					80
204	Arg	Lys	Lys	Ser	Leu	Met	Ile	Gly	Ala	Ile	Leu	Phe	Val	Ala	Gly	Ser
205		_			85					90	•				95	
207	Leu	Phe	Ser	Ala	Ala	Ala	Pro	Asn	Val	Glu	Val	Leu	Ile	Leu	Ser	Arg
208				100					105					110		
210	Val	Leu	Leu	Gly	Leu	Ala	Val	Gly	Val	Ala	Ser	Tyr	Thr	Ala	Pro	Leu
211			115					120					125			
213	Tyr	Leu	Ser	Glu	Ile	Ala	Pro	Glu	Lys	Ile	Arg	Gly	Ser	Met	Ile	Ser
214		130					135					140				
216	Met	Tyr	Gln	Leu	Met	Ile	Thr	Ile	Gly	Ile	Leu	Gly	Ala	Tyr	Leu	Ser
217	145	_		•		150			•		155					160
219	Asp	Thr	Ala	Phe	Ser	Tyr	Thr	Gly	Ala	Trp	Arg	Trp	Met	Leu	Gly	Val
220					165					170					175	
222	Ile	Ile	Ile	Pro	Ala	Ile	Leu	Leu	Leu	Ile	Gly	Val	Phe	Phe	Leu	Pro
223				180					185			•		190		
225	Asp	Ser	Pro	Arg	Trp	Phe	Ala	Ala	Lys	Arg	Arg	Phe	Val	Asp	Ala	Glu
226			195					200					205			
228	Arg	Val	Leu	Leu	Arg	Leu	Arg	Asp	Thr	Ser	Ala	Glu	Ala	Lys	Arg	Glu
229	_	210					215					220				
231	Leu	Asp	Glu	Ile	Arg	Glu	Ser	Leu	Gln	Val	Lys	Gln	Ser	Gly	Trp	Ala
232	225					230					235					240
234	Leu	Phe	Lys	Glu	Asn	Ser	Asn	Phe	Arg	Arg	Ala	Val	Phe	Leu	Gly	Val
235					245					250			•		255	
237	Leu	Leu	Gln	Val	Met	Gln	Gln	Phe	Thr	Gly	Met	Asn	Val	Ile	Met	Tyr
238				260					265					270		
241	Tyr	Ala	Pro	Lys	Ile	Phe	Glu	Leu	Ala	Gly	Tyr	Thr	Asn	Thr	Thr	Glu
242			275					280					285			
244	Gln	Met	Trp	Gly	Thr	Val	Ile	Val	Gly	Leu	Thr	Asn	Val	Leu	Ala	Thr
245		290					295					300				
247	Phe	Ile	Ala	Ile	Gly	Leu	Val	Asp	Arg	Trp	Gly	Arg	Lys	Pro	Thr	Leu
248	305	•				310					315					320
250	Thr	Leu	Gly	Phe	Leu	Val	Met	Ala	Ala	Gly	Met	Gly	Val	Leu	Gly	Thr
251					325					330					335	•
253	Met	Met	His	Ile	Gly	Ile	His	Ser	Pro	Ser	Ala	Gln	Tyr		Ala	Ile
254		•		340	•		· .•		345					350		
256	Ala	Met	Leu	Leu	Met	Phe	Ile	Val	Gly	Phe	Ala	Met	Ser	Ala	Gly	Pro
257	•		355					360					365			
259	Leu	Ile	Trp	Val	Leu	Cys	Ser	Glu	Ile	Gln	Pro	Leu	Lys	Gly	Arg	Asp
260		370					375					380				
262	Phe	Gly	Ile	Thr	Cys	Ser	Thr	Ala	Thr	Asn	Trp	Ile	Ala	Asn	Met	Ile
263	385					390					395					400
265	Val	Gly	Ala	Thr	Phe	Leu	Thr	Met	Leu	Asn	Thr	Leu	Gly	Asn		Asn
266		_			405					410					415	
268	Thr	Phe	Trp	Val	Tyr	Ala	Ala	Leu	Asn	Val	Leu	Phe	Ile		Leu	Thr
269				420					425				,	430		

1 age J Ut /

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/812,315

DATE: 04/06/2004 TIME: 10:51:23

Input Set : A:\Sequence Listing 81000.txt
Output Set: N:\CRF4\04062004\J812315.raw

271 Leu Trp Leu Val Pro Glu Thr Lys His Val Ser Leu Glu His Ile Glu

272 435 440 445

274 Arg Asn Leu Met Lys Gly Arg Lys Leu Arg Glu Ile Gly Ala His Asp

75 450 455 460

1 agv v vi /

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/812,315

DATE: 04/06/2004

TIME: 10:51:24

Input Set : A:\Sequence Listing 81000.txt Output Set: N:\CRF4\04062004\J812315.raw

:11 M:270 C: Current Application Number differs, Replaced Current Application No

:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

:22 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 :37 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2 :52 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3 :61 M:283 W: Missing Blank Line separator, <400> field identifier